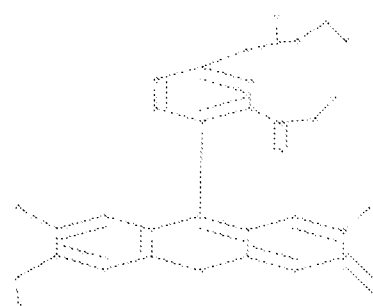
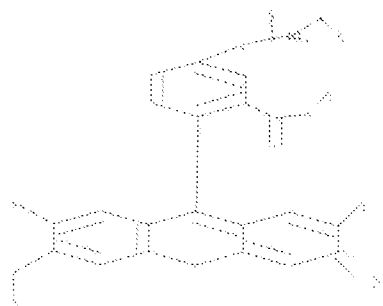


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=>

Uploading C:\Program Files\Stnexp\Queries\11767334.etr



chain nodes :

21 22 23 24 25 27 29 30 33 35 36 37 38 39 40

ring nodes :

1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20

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chain bonds :

2-22 3-30 7-15 12-29 13-21 18-35 20-23 22-33 23-24 23-25 25-27  
35-36 36-37 36-38 38-39 39-40

ring bonds :

1-2 1-6 2-3 3-4 4-5 5-6 5-7 6-10 7-8 8-9 8-11 9-10 9-14 11-12  
12-13 13-14 15-16 15-20 16-17 17-18 18-19 19-20

exact/norm bonds :

2-22 3-30 12-29 13-21 18-35 22-33 23-24 23-25 25-27 35-36 36-37  
39-40

exact bonds :

5-7 6-10 7-8 7-15 8-9 8-11 9-10 9-14 11-12 12-13 13-14 20-23  
36-38 38-39

normalized bonds :

1-2 1-6 2-3 3-4 4-5 5-6 15-16 15-20 16-17 17-18 18-19 19-20

isolated ring systems :

containing 1 : 15 :

G1:H,Ak

G2:H,K

G3:H,Ak,C

G4:H,H<sub>y</sub>

Match level :

1:Atom 2:Atom 3:Atom 4:Atom 5:Atom 6:Atom 7:Atom 8:Atom 9:Atom  
10:Atom 11:Atom 12:Atom 13:Atom 14:Atom 15:Atom 16:Atom 17:Atom  
18:Atom 19:Atom 20:Atom 21:CLASS 22:CLASS 23:CLASS 24:CLASS 25:CLASS  
27:CLASS 29:CLASS 30:CLASS 33:CLASS 35:CLASS 36:CLASS 37:CLASS  
38:CLASS 39:CLASS 40:CLASS

L1 STRUCTURE UPLOADED

=> s 11

SAMPLE SEARCH INITIATED 19:37:36 FILE 'REGISTRY'

SAMPLE SCREEN SEARCH COMPLETED - 8 TO ITERATE

100.0% PROCESSED 8 ITERATIONS

0 ANSWERS

SEARCH TIME: 00.00.01

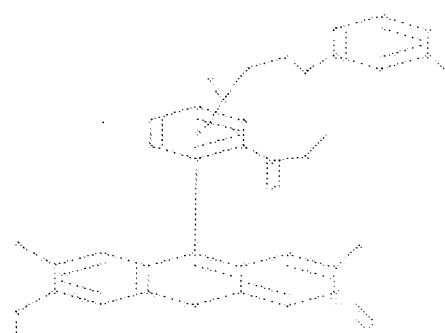
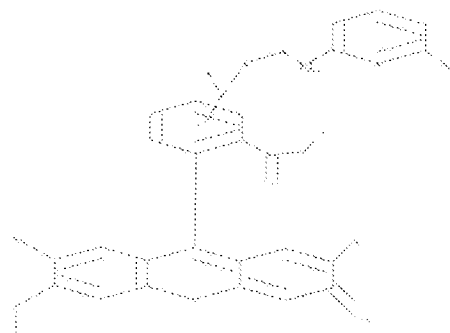
FULL FILE PROJECTIONS: ONLINE \*\*COMPLETE\*\*

BATCH \*\*COMPLETE\*\*

PROJECTED ITERATIONS: 8 TO 329

PROJECTED ANSWERS: 0 TO 0

L2 0 SEA SSS SAM L1



chain nodes :

21 22 23 24 25 27 29 30 33 36 37 38 39 40 47 50

ring nodes :

1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 41  
42 43 44 45 46

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chain bonds :

2-22 3-30 7-15 12-29 13-21 20-23 22-33 23-24 23-25 25-27 36-37  
36-38 36-50 39-40 39-50 40-42 46-47

ring bonds :

1-2 1-6 2-3 3-4 4-5 5-6 5-7 6-10 7-8 8-9 8-11 9-10 9-14 11-12  
12-13 13-14 15-16 15-20 16-17 17-18 18-19 19-20 41-42 41-46 42-43  
43-44 44-45 45-46

exact/norm bonds :

2-22 3-30 12-29 13-21 22-33 23-24 23-25 25-27 36-37 36-50 39-50  
46-47

exact bonds :

5-7 6-10 7-8 7-15 8-9 8-11 9-10 9-14 11-12 12-13 13-14 20-23  
36-38 39-40 40-42

normalized bonds :

1-2 1-6 2-3 3-4 4-5 5-6 15-16 15-20 16-17 17-18 18-19 19-20  
41-42 41-46 42-43 43-44 44-45 45-46

isolated ring systems :

containing 1 : 15 : 41 :

G1:N,Ak

G2:H,X

G3:H,Ak,C

G4:H,H<sub>y</sub>

Match level :

1:Atom 2:Atom 3:Atom 4:Atom 5:Atom 6:Atom 7:Atom 8:Atom 9:Atom  
10:Atom 11:Atom 12:Atom 13:Atom 14:Atom 15:Atom 16:Atom 17:Atom  
18:Atom 19:Atom 20:Atom 21:CLASS 22:CLASS 23:CLASS 24:CLASS 25:CLASS  
27:CLASS 29:CLASS 30:CLASS 33:CLASS 36:CLASS 37:CLASS 38:CLASS  
39:CLASS 40:CLASS 41:CLASS 42:CLASS 43:Atom 44:Atom 45:Atom 46:Atom  
47:CLASS 50:CLASS 51:Atom

L4 STRUCTURE UPLOADED

=> s 14

SAMPLE SEARCH INITIATED 19:44:35 FILE 'REGISTRY'

SAMPLE SCREEN SEARCH COMPLETED - 0 TO ITERATE

100.0% PROCESSED 0 ITERATIONS

0 ANSWERS

SEARCH TIME: 00.00.01

FULL FILE PROJECTIONS: ONLINE \*\*COMPLETE\*\*  
BATCH \*\*COMPLETE\*\*

PROJECTED ITERATIONS: 0 TO 0

PROJECTED ANSWERS: 0 TO 0